

# SEQUENCE LISTING

<110> Ruben et al.

<120> Cytokine Receptor Common Gamma Chain Like

<130> PF466P2

<150> 60/269,876

<151> 2001-02-21

<150> PCT/US00/22493

<151> 2000-08-17

<150> 09/376,430

<151> 1999-08-18

<150> 09/263,626

<151> 1999-03-05

<150> PCT/US99/05068

<151> 1999-03-05

<150> 60/086,505

<151> 1998-05-22

<150> 60/078,563

<151> 1998-03-19

<160> 32

<170> PatentIn Ver. 2.1

<210> 1

<211> 1573

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (13)..(1125)

<400> 1

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Met Gly Arg Leu Val Leu Leu Trp Gly Ala Ala Val Phe

1

5

10

ctg ctg gga ggc tgg atg gct ttg ggg caa gga gga gca gca gaa gga 99

Leu Leu Gly Gly Trp Met Ala Leu Gly Gln Gly Gly Ala Ala Glu Gly

15

20

25

gta cag att cag atc atc tac ttc aat tta gaa acc gtg cag gtg aca 147

Val Gln Ile Gln Ile Ile Tyr Phe Asn Leu Glu Thr Val Gln Val Thr

30

35

40

45

tgg aat gcc agc aaa tac tcc agg acc aac ctg act ttc cac tac aga 195

Trp Asn Ala Ser Lys Tyr Ser Arg Thr Asn Leu Thr Phe His Tyr Arg

50

55

60

ttc aac ggt gat gag gcc tat gac cag tgc acc aac tac ctt ctc cag	243
Phe Asn Gly Asp Glu Ala Tyr Asp Gln Cys Thr Asn Tyr Leu Leu Gln	
65 70 75	
gaa ggt cac act tcg ggg tgc ctc cta gac gca gag cag cga gac gac	291
Glu Gly His Thr Ser Gly Cys Leu Leu Asp Ala Glu Gln Arg Asp Asp	
80 85 90	
att ctc tat ttc tcc atc agg aat ggg acg cac ccc gtt ttc acc gca	339
Ile Leu Tyr Phe Ser Ile Arg Asn Gly Thr His Pro Val Phe Thr Ala	
95 100 105	
agt cgc tgg atg gtt tat tac ctg aaa ccc agt tcc ccg aag cac gtg	387
Ser Arg Trp Met Val Tyr Tyr Leu Lys Pro Ser Ser Pro Lys His Val	
110 115 120 125	
aga ttt tcg tgg cat cag gat gca gtg acg gtg acg tgt tct gac ctg	435
Arg Phe Ser Trp His Gln Asp Ala Val Thr Val Thr Cys Ser Asp Leu	
130 135 140	
tcc tac ggg gat ctc ctc tat gag gtt cag tac cgg agc ccc ttc gac	483
Ser Tyr Gly Asp Leu Leu Tyr Glu Val Gln Tyr Arg Ser Pro Phe Asp	
145 150 155	
acc gag tgg cag tcc aaa cag gaa aat acc tgc aac gtc acc ata gaa	531
Thr Glu Trp Gln Ser Lys Gln Glu Asn Thr Cys Asn Val Thr Ile Glu	
160 165 170	
ggc ttg gat gcc gag aag tgt tac tct ttc tgg gtc agg gtg aag gct	579
Gly Leu Asp Ala Glu Lys Cys Tyr Ser Phe Trp Val Arg Val Lys Ala	
175 180 185	
atg gag gat gta tat ggg cca gac aca tac cca agc gac tgg tca gag	627
Met Glu Asp Val Tyr Gly Pro Asp Thr Tyr Pro Ser Asp Trp Ser Glu	
190 195 200 205	
gtg aca tgc tgg cag aga ggc gag att cgg gat gcc tgt gca gag aca	675
Val Thr Cys Trp Gln Arg Gly Glu Ile Arg Asp Ala Cys Ala Glu Thr	
210 215 220	
cca acg cct ccc aaa cca aag ctg tcc aaa ttt att tta att tcc agc	723
Pro Thr Pro Pro Lys Pro Lys Leu Ser Lys Phe Ile Leu Ile Ser Ser	
225 230 235	
ctg gcc atc ctt ctg atg gtg tct ctc ctc ctt ctg tct tta tgg aaa	771
Leu Ala Ile Leu Leu Met Val Ser Leu Leu Leu Leu Ser Leu Trp Lys	
240 245 250	
tta tgg aga gtg aag aag ttt ctc att ccc agc gtg cca gac ccg aaa	819
Leu Trp Arg Val Lys Lys Phe Leu Ile Pro Ser Val Pro Asp Pro Lys	
255 260 265	
tcc atc ttc ccc ggg ctc ttt gag ata cac caa ggg aac ttc cag gag	867
Ser Ile Phe Pro Gly Leu Phe Glu Ile His Gln Gly Asn Phe Gln Glu	
270 275 280 285	
tgg atc aca gac acc cag aac gtg gcc cac ctc cac aag atg gca ggt	915
Trp Ile Thr Asp Thr Gln Asn Val Ala His Leu His Lys Met Ala Gly	
290 295 300	

gca gag caa gaa agt ggc ccc gag gag ccc ctg gta gtc cag ttg gcc 963  
 Ala Glu Gln Glu Ser Gly Pro Glu Glu Pro Leu Val Val Gln Leu Ala  
 305 310 315  
 aag act gaa gcc gag tct ccc agg atg ctg gac cca cag acc gag gag 1011  
 Lys Thr Glu Ala Glu Ser Pro Arg Met Leu Asp Pro Gln Thr Glu Glu  
 320 325 330  
 aaa gag gcc tct ggg gga tcc ctc cag ctt ccc cac cag ccc ctc caa 1059  
 Lys Glu Ala Ser Gly Gly Ser Leu Gln Leu Pro His Gln Pro Leu Gln  
 335 340 345  
 ggc ggt gat gtg gtc aca atc ggg ggc ttc acc ttt gtg atg aat gac 1107  
 Gly Gly Asp Val Val Thr Ile Gly Gly Phe Thr Phe Val Met Asn Asp  
 350 355 360 365  
 cgc tcc tac gtg gcg ttg tgatggacac accactgtca aagtcaacgt 1155  
 Arg Ser Tyr Val Ala Leu  
 370  
 caggatccac gttgacattt aaagacagag gggactgtcc cggggactcc acaccaccat 1215  
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 <212> PRT  
 <213> Homo sapiens

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 Gly Trp Met Ala Leu Gly Gln Gly Gly Ala Ala Glu Gly Val Gln Ile  
 20 25 30  
 Gln Ile Ile Tyr Phe Asn Leu Glu Thr Val Gln Val Thr Trp Asn Ala  
 35 40 45  
 Ser Lys Tyr Ser Arg Thr Asn Leu Thr Phe His Tyr Arg Phe Asn Gly  
 50 55 60  
 Asp Glu Ala Tyr Asp Gln Cys Thr Asn Tyr Leu Leu Gln Glu Gly His  
 65 70 75 80  
 Thr Ser Gly Cys Leu Leu Asp Ala Glu Gln Arg Asp Asp Ile Leu Tyr  
 85 90 95

Phe Ser Ile Arg Asn Gly Thr His Pro Val Phe Thr Ala Ser Arg Trp  
 100 105 110  
 Met Val Tyr Tyr Leu Lys Pro Ser Ser Pro Lys His Val Arg Phe Ser  
 115 120 125  
 Trp His Gln Asp Ala Val Thr Val Thr Cys Ser Asp Leu Ser Tyr Gly  
 130 135 140  
 Asp Leu Leu Tyr Glu Val Gln Tyr Arg Ser Pro Phe Asp Thr Glu Trp  
 145 150 155 160  
 Gln Ser Lys Gln Glu Asn Thr Cys Asn Val Thr Ile Glu Gly Leu Asp  
 165 170 175  
 Ala Glu Lys Cys Tyr Ser Phe Trp Val Arg Val Lys Ala Met Glu Asp  
 180 185 190  
 Val Tyr Gly Pro Asp Thr Tyr Pro Ser Asp Trp Ser Glu Val Thr Cys  
 195 200 205  
 Trp Gln Arg Gly Glu Ile Arg Asp Ala Cys Ala Glu Thr Pro Thr Pro  
 210 215 220  
 Pro Lys Pro Lys Leu Ser Lys Phe Ile Leu Ile Ser Ser Leu Ala Ile  
 225 230 235 240  
 Leu Leu Met Val Ser Leu Leu Leu Leu Ser Leu Trp Lys Leu Trp Arg  
 245 250 255  
 Val Lys Lys Phe Leu Ile Pro Ser Val Pro Asp Pro Lys Ser Ile Phe  
 260 265 270  
 Pro Gly Leu Phe Glu Ile His Gln Gly Asn Phe Gln Glu Trp Ile Thr  
 275 280 285  
 Asp Thr Gln Asn Val Ala His Leu His Lys Met Ala Gly Ala Glu Gln  
 290 295 300  
 Glu Ser Gly Pro Glu Glu Pro Leu Val Val Gln Leu Ala Lys Thr Glu  
 305 310 315 320  
 Ala Glu Ser Pro Arg Met Leu Asp Pro Gln Thr Glu Glu Lys Glu Ala  
 325 330 335  
 Ser Gly Gly Ser Leu Gln Leu Pro His Gln Pro Leu Gln Gly Gly Asp  
 340 345 350  
 Val Val Thr Ile Gly Gly Phe Thr Phe Val Met Asn Asp Arg Ser Tyr  
 355 360 365  
 Val Ala Leu  
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<210> 3

<211> 379

<212> PRT

<213> Homo sapiens

<400> 3

Met Leu Lys Pro Pro Leu Pro Leu Arg Ser Leu Leu Phe Leu Gln Leu  
1 5 10 15

Pro Leu Leu Gly Val Gly Leu Asn Pro Lys Phe Leu Thr Pro Ser Gly  
20 25 30

Asn Glu Asp Ile Gly Gly Lys Pro Gly Thr Gly Gly Asp Phe Phe Leu  
35 40 45

Thr Ser Thr Pro Ala Gly Thr Leu Asp Val Ser Thr Leu Pro Leu Pro  
50 55 60

Lys Val Gln Cys Phe Val Phe Asn Val Glu Tyr Met Asn Cys Thr Trp  
65 70 75 80

Asn Ser Ser Ser Glu Pro Gln Pro Asn Asn Leu Thr Leu His Tyr Gly  
85 90 95

Tyr Arg Asn Phe Asn Gly Asp Asp Lys Leu Gln Glu Cys Gly His Tyr  
100 105 110

Leu Phe Ser Glu Gly Ile Thr Ser Gly Cys Trp Phe Gly Lys Lys Glu  
115 120 125

Ile Arg Leu Tyr Glu Thr Phe Val Val Gln Leu Gln Asp Pro Arg Glu  
130 135 140

His Arg Lys Gln Pro Lys Gln Met Leu Lys Leu Gln Asp Leu Val Ile  
145 150 155 160

Pro Trp Ala Pro Glu Asn Leu Thr Leu Arg Asn Leu Ser Glu Phe Gln  
165 170 175

Leu Glu Leu Ser Trp Ser Asn Arg Tyr Leu Asp His Cys Leu Glu His  
180 185 190

Leu Val Gln Tyr Arg Ser Asp Arg Asp Arg Ser Trp Thr Glu Gln Ser  
195 200 205

Val Asp His Arg His Ser Phe Ser Leu Pro Ser Val Asp Ala Gln Lys  
210 215 220

Leu Tyr Thr Phe Arg Val Arg Ser Arg Tyr Asn Pro Leu Cys Gly Ser  
225 230 235 240

Ala Gln His Trp Ser Asp Trp Ser Tyr Pro Ile His Trp Gly Ser Asn  
245 250 255

Thr Ser Lys Glu Asn Ile Glu Asn Pro Glu Asn Pro Ser Leu Phe Ala  
260 265 270

Leu Glu Ala Val Leu Ile Pro Leu Gly Ser Met Gly Leu Ile Val Ser  
275 280 285

Leu Ile Cys Val Tyr Cys Trp Leu Glu Arg Thr Met Pro Arg Ile Pro

290	295	300
Thr Leu Lys Asn Leu Glu Asp Leu Val Thr Glu Tyr Gln Gly Asn Phe		
305	310	315 320
Ser Ala Trp Ser Gly Val Ser Lys Gly Leu Ala Glu Ser Leu Gln Pro		
	325	330 335
Asp Tyr Ser Glu Arg Leu Cys His Val Ser Glu Ile Pro Pro Lys Gly		
	340	345 350
Gly Glu Gly Pro Gly Gly Ser Pro Cys Ser Gln His Ser Pro Tyr Trp		
	355	360 365
Ala Pro Pro Cys Tyr Thr Leu Lys Pro Glu Pro		
	370	375

<210> 4  
 <211> 733  
 <212> DNA  
 <213> Homo sapiens

<400> 4  
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 aattcgaggg tgcaccgtca gtcttctctt tccccccaaa acccaaggac accctcatga 120  
 tctcccggac tcctgaggtc acatgcgtgg tggtaggacgt aagccacgaa gaccctgagg 180  
 tcaagttcaa ctggtacgtg gacggcgtgg aggtgcataa tgccaagaca aagccgcggg 240  
 aggagcagta caacagcacg taccgtgtgg tcagcgtcct caccgtcctg caccaggact 300  
 ggctgaatgg caaggagtac aagtgcagg tctccaacaa agccctccca acccccatcg 360  
 agaaaaccat ctccaaagcc aaagggcagc cccgagaacc acaggtgtac accctgcccc 420  
 catcccggga tgagctgacc aagaaccagg tcagcctgac ctgcctgggtc aaaggcttct 480  
 atccaagcga catcgccgtg gagtgggaga gcaatgggca gccggagaa aactacaaga 540  
 ccacgcctcc cgtgctggac tccgacggct ccttcttctt ctacagcaag ctcaccgtgg 600  
 acaagagcag gtggcagcag gggaacgtct tctcatgctc cgtgatgcac gaggtctctg 660  
 acaaccacta cagcagaag agcctctccc tgtctccggg taaatgagtg cgacggccgc 720  
 gactctagag gat 733

<210> 5  
 <211> 5  
 <212> PRT  
 <213> Homo sapiens

<220>  
 <221> SITE  
 <222> (3)  
 <223> Xaa equals any amino acid

<400> 5  
 Trp Ser Xaa Trp Ser  
 1 5

<210> 6  
 <211> 86  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <221> Primer\_Bind  
 <223> Synthetic sequence with 4 tandem copies of the GAS binding site found in the IRF1 promoter (Rothman et al., Immunity 1:457-468 (1994)), 18 nucleotides complementary to the SV40 early promoter, and a Xho I restriction site.

<400> 6  
 gcgcctcgag atttccccga aatctagatt tccccgaaat gatttccccg aaatgatttc 60  
 ccgaaatat ctgccatctc aattag 86

<210> 7  
 <211> 27  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <221> Primer\_Bind  
 <223> Synthetic sequence complementary to the SV40 promoter; includes a Hind III restriction site.

<400> 7  
 gcggcaagct ttttgcaaag cctaggc 27

<210> 8  
 <211> 271  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <221> Protein\_Bind  
 <223> Synthetic promoter for use in biological assays; includes GAS binding sites found in the IRF1 promoter (Rothman et al., Immunity 1:457-468 (1994)).

<400> 8  
 ctcgagattt ccccgaaatc tagatttccc cgaaatgatt tccccgaaat gatttccccg 60  
 aaatatctgc catctcaatt agtcagcaac catagtcccc cccctaactc cgcccatccc 120  
 gccctaact ccgcccagtt ccgcccattc tccgccccat ggctgactaa ttttttttat 180  
 ttatgcagag gccgaggccg cctcggcctc tgagctattc cagaagtagt gaggaggctt 240  
 ttttgagggc ctaggctttt gcaaaaagct t 271

<210> 9  
 <211> 32  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <221> Primer\_Bind  
 <223> Synthetic primer complementary to human genomic EGR-1 promoter sequence (Sakamoto et al., Oncogene 6:867-871 (1991)); including an Xho I restriction site.

<400> 9  
 gcgctcgagg gatgacagcg atagaacccc gg 32

<210> 10  
 <211> 31  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <221> Primer\_Bind  
 <223> Synthetic primer complementary to human genomic EGR-1 promoter sequence (Sakamoto et al., Oncogene 6:867-871 (1991)); including an Hind III restriction site.  
  
 <400> 10  
 gcgaagcttc gcgactcccc ggatccgcct c 31  
  
 <210> 11  
 <211> 12  
 <212> DNA  
 <213> Homo sapiens  
  
 <400> 11  
 ggggactttc cc 12  
  
 <210> 12  
 <211> 73  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <221> Primer\_Bind  
 <223> Synthetic primer with 4 tandem copies of the NF-KB binding site (GGGGACTTTCCC), 18 nucleotides complementary to the 5' end of the SV40 early promoter sequence, and a XhoI restriction site.  
  
 <400> 12  
 gcggcctcga ggggactttc ccggggactt tccggggact ttccgggact ttccatcctg 60  
 ccattctcaat tag 73  
  
 <210> 13  
 <211> 256  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <221> Protein\_Bind  
 <223> Synthetic promoter for use in biological assays; including NF-KB binding sites.  
  
 <400> 13  
 ctcgagggga ctttcccggg gactttccgg ggactttccg ggactttcca tctgccatct 60  
 caattagtca gcaaccatag tcccgccct aactccgcc atcccgcgcc taactccgcc 120  
 cagttccgcc cattctccgc cccatggctg actaattttt ttattttatg cagaggccga 180  
 ggccgctcgc gcctctgagc tattccagaa gtagtgagga ggcttttttg gaggcctagg 240  
 cttttgcaaa aagctt 256



<210> 14  
<211> 29  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> Primer\_Bind  
<223> Synthetic primer containing the Nco I restriction site followed a number of nucleotides of the amino terminal coding sequence of CRCGCL

<400> 14  
gttaggcat gggaggagca gcagaagga 29

<210> 15  
<211> 33  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> Primer\_Bind  
<223> Synthetic primer containing the BglII restriction site followed by a number nucleotides complementary to the 3' end of the coding sequence of CRCGCL

<400> 15  
ggtaaagat ctcaacgccca cgtaggagcg gtc 33

<210> 16  
<211> 38  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> Primer\_Bind  
<223> Synthetic primer to amplify the cDNA sequence encoding the full length CRCGCL protein including the AUG initiation codon and the naturally associated leader,also contains a BglII site,and Kozak signal

<400> 16  
ccggttagat ctgccatcat ggctttgggg caaggagg 38

<210> 17  
<211> 36  
<212> DNA  
<213> Artificial Sequence

<220>  
<221> Primer\_Bind  
<223> Synthetic primercontaining the XbaI restriction site followed by a number of nucleotides complementary to the 3' noncoding sequence of SEQ ID NO:1

<400> 17  
ccggtttcta gatcacaagg ccacgtagga gcggtc 36

<210> 18  
<211> 7  
<212> PRT  
<213> Homo sapiens

<220>  
<221> SITE  
<222> (1)  
<223> Xaa equals Ser, Thr, Gly or Leu

<220>  
<221> SITE  
<222> (2)  
<223> Xaa equals any amino acid

<220>  
<221> SITE  
<222> (4)  
<223> Xaa equals Ser or Gly

<220>  
<221> SITE  
<222> (5)  
<223> Xaa equals any amino acid

<400> 18  
Xaa Xaa Trp Xaa Xaa Trp Ser  
1 5

<210> 19  
<211> 7  
<212> PRT  
<213> Homo sapiens

<220>  
<221> SITE  
<222> (2)  
<223> Xaa equals any amino acid

<220>  
<221> SITE  
<222> (5)  
<223> Xaa equals any amino acid

<400> 19  
Thr Xaa Pro Ser Xaa Trp Ser  
1 5

<210> 20  
<211> 7  
<212> PRT  
<213> Homo sapiens

<220>  
<221> SITE

<222> (2)  
 <223> Xaa equals Pro or Glu  
  
 <220>  
 <221> SITE  
 <222> (3)  
 <223> Xaa equals any amino acid  
  
 <220>  
 <221> SITE  
 <222> (4)  
 <223> Xaa equals Val or Ile  
  
 <220>  
 <221> SITE  
 <222> (6)  
 <223> Xaa equals Asn, Ser or Asp

<400> 20  
 Trp Xaa Xaa Xaa Pro Xaa Pro  
     1                    5

<210> 21  
 <211> 7  
 <212> PRT  
 <213> Homo sapiens

<220>  
 <221> SITE  
 <222> (3)  
 <223> Xaa equals any amino acid

<400> 21  
 Ile Pro Xaa Val Pro Asp Pro  
     1                    5

<210> 22  
 <211> 54  
 <212> PRT  
 <213> Homo sapiens

<400> 22  
 Gln Ile Gln Ile Ile Tyr Phe Asn Leu Glu Thr Val Gln Val Thr Trp  
     1                    5                    10                    15  
  
 Asn Ala Ser Lys Tyr Ser Arg Thr Asn Leu Thr Phe His Tyr Arg Phe  
                     20                    25                    30  
  
 Asn Gly Asp Glu Ala Tyr Asp Gln Cys Thr Asn Tyr Leu Leu Gln Glu  
                     35                    40                    45  
  
 Gly His Thr Ser Gly Cys  
     50

<210> 23  
 <211> 30

<212> PRT  
<213> Homo sapiens

<400> 23

Arg Arg His Ser Leu Phe Leu His Gln Glu Trp Asp Ala Pro Arg Phe  
1 5 10 15

His Arg Lys Ser Leu Asp Gly Leu Leu Pro Glu Thr Gln Phe  
20 25 30

<210> 24

<211> 81

<212> PRT

<213> Homo sapiens

<400> 24

Leu Leu Tyr Glu Val Gln Tyr Arg Ser Pro Phe Asp Thr Glu Trp Gln  
1 5 10 15

Ser Lys Gln Glu Asn Thr Cys Asn Val Thr Ile Glu Gly Leu Asp Ala  
20 25 30

Glu Lys Cys Tyr Ser Phe Trp Val Arg Val Lys Ala Met Glu Asp Val  
35 40 45

Tyr Gly Pro Asp Thr Tyr Pro Ser Asp Trp Ser Glu Val Thr Cys Trp  
50 55 60

Gln Arg Gly Glu Ile Arg Asp Ala Cys Ala Glu Thr Pro Thr Pro Pro  
65 70 75 80

Lys

<210> 25

<211> 181

<212> PRT

<213> Homo sapiens

<220>

<221> SITE

<222> (68)

<223> Xaa equals any amino acid

<220>

<221> SITE

<222> (73)

<223> Xaa equals any amino acid

<220>

<221> SITE

<222> (88)

<223> Xaa equals any amino acid

<400> 25

Met Glu Asp Val Tyr Gly Pro Asp Thr Tyr Pro Ser Asp Trp Ser Glu  
1 5 10 15

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			20					25					30		
Pro	Thr	Pro	Pro	Lys	Pro	Lys	Leu	Ser	Lys	Phe	Ile	Leu	Ile	Ser	Ser
		35					40					45			
Leu	Ala	Ile	Leu	Leu	Met	Val	Ser	Leu	Leu	Leu	Leu	Ser	Leu	Trp	Lys
		50				55					60				
Leu	Trp	Arg	Xaa	Lys	Lys	Phe	Leu	Xaa	Pro	Ser	Val	Pro	Asp	Pro	Lys
	65				70					75					80
Ser	Ile	Phe	Pro	Gly	Leu	Phe	Xaa	Ile	His	Gln	Gly	Asn	Phe	Gln	Glu
				85					90					95	
Trp	Ile	Thr	Asp	Thr	Gln	Asn	Val	Ala	His	Leu	His	Lys	Met	Ala	Gly
			100					105					110		
Ala	Glu	Gln	Glu	Ser	Gly	Pro	Glu	Glu	Pro	Leu	Val	Val	Gln	Leu	Ala
		115					120					125			
Lys	Thr	Glu	Ala	Glu	Ser	Pro	Arg	Met	Leu	Asp	Pro	Gln	Thr	Glu	Glu
	130					135					140				
Lys	Glu	Ala	Ser	Gly	Gly	Ser	Leu	Gln	Leu	Pro	His	Gln	Pro	Leu	Gln
	145				150					155					160
Gly	Gly	Asp	Val	Val	Thr	Ile	Gly	Gly	Phe	Thr	Phe	Val	Met	Asn	Asp
			165					170						175	
Arg	Ser	Tyr	Val	Ala											
			180												

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<210> 26
<211> 1567
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (830)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (416)
<223> y equals c or t

<220>
<221> misc_feature
<222> (784)
<223> m equals a or c

<220>
<221> misc_feature
<222> (785)
<223> y equals c or t

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<400> 26
gggcatgggg cggctgggttc tgetgtgggg agctgccgtc tttctgctgg gaggctggat 60
ggctttgggg caaggaggag cagcagaagg agtacagatt caratcatct acttcaattt 120
agaaaccgtg caggtgacat ggaatgccag caaatactcc aggaccaacc tgactttcca 180
ctacagattc aacgggtgatg aggocctatga ccagtgcacc aactaccttc tccaggaagg 240
tcacacttcg gggtgccctcc tagacgcasa gcagcgagac gacatttctct atttctccat 300
caggaatggg acgcaccccg ttttcaccgc aagtcgctgg atggtttatt acctgaaacc 360
cagttccccg aagcacgtga gatttcgtgg catcaggaaw gacggtgacg tgttcycgac 420
ctgtcctacg gggatctcct ctatgaggtt cagtaccgga gccccttcga caccgagtgg 480
cagtccaaac aggaaaatac ctgcaacgtc accatagaag gcttgatgc cgagaagtgt 540
tactctttct gggtcagggt gaaggctatg gaggatgtat atggggccaga cacataccca 600
agcgactggg cagagggtgac atgctggcag agaggcgaga ttcgggatgc ctgtgcagag 660
acaccaacgc ctcccaaacc aaagctgtcc aaattttatt taatttccag cctggccatc 720
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<210> 27
<211> 170
<212> PRT
<213> Homo sapiens

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<220>
<221> SITE
<222> (89)
<223> Xaa equals any amino acid

```

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<220>
<221> SITE
<222> (132)
<223> Xaa equals any amino acid

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<220>
<221> SITE
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<223> Xaa equals any amino acid

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<400> 27
Met Gly Arg Leu Val Leu Leu Trp Gly Ala Ala Val Phe Leu Leu Gly
 1           5           10           15

Gly Trp Met Ala Leu Gly Gln Gly Gly Ala Ala Glu Gly Val Gln Ile
          20           25           30

Gln Ile Ile Tyr Phe Asn Leu Glu Thr Val Gln Val Thr Trp Asn Ala

```

35	40	45
Ser Lys Tyr Ser Arg Thr Asn Leu Thr Phe His Tyr Arg Phe Asn Gly		
50	55	60
Asp Glu Ala Tyr Asp Gln Cys Thr Asn Tyr Leu Leu Gln Glu Gly His		
65	70	75
Thr Ser Gly Cys Leu Leu Asp Ala Xaa Gln Arg Asp Asp Ile Leu Tyr		
	85	90
Phe Ser Ile Arg Asn Gly Thr His Pro Val Phe Thr Ala Ser Arg Trp		
	100	105
Met Val Tyr Tyr Leu Lys Pro Ser Ser Pro Lys His Val Arg Phe Arg		
	115	120
Gly Ile Arg Xaa Asp Gly Asp Val Phe Xaa Thr Cys Pro Thr Gly Ile		
	130	135
Ser Ser Met Arg Phe Ser Thr Gly Ala Pro Ser Thr Pro Ser Gly Ser		
145	150	155
Pro Asn Arg Lys Ile Pro Ala Thr Ser Pro		
	165	170

<210> 28  
 <211> 36  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <221> Primer\_Bind  
 <223> Synthetic primer to amplify the cDNA sequence encoding the amino acids Met 1 to Lys 231 including the AUG initiation codon and the naturally associated leader,also contains a BglII site,and Kozak signal

<400> 28  
 ccggttagat ctgccatcat ggggcggctg gttctg 36

<210> 29  
 <211> 31  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <221> Primer\_Bind  
 <223> Synthetic primer to amplify the cDNA sequence encoding the amino acids Met 1 to Lys 231,also contains a BglII site

<400> 29  
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<210> 30  
 <211> 4  
 <212> PRT

<213> Homo sapiens

<220>

<221> SITE

<222> (2)

<223> Xaa equals any amino acid

<400> 30

Trp Xaa Trp Ser

1

<210> 31

<211> 34

<212> DNA

<213> Artificial Sequence

<220>

<221> Primer\_Bind

<223> Synthetic primer to amplify the cDNA sequence encoding the amino acids Met 1 to Lys 231, also contains a XbaI site

<400> 31

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34

<210> 32

<211> 144

<212> DNA

<213> Artificial Sequence

<220>

<221> Primer\_Bind

<223> Synthetic primer to amplify DNA encoding amino acids M1-K231 of SEQ ID NO:2, contains a BglII restriction site.

<400> 32

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gccatcatgg ggcggctggt tctg 144